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### RECEIVED MAY 1 4 1987

## ecology and environment, inc.

111 WEST JACKSON BLVD., CHICAGO	, ILLINOIS 60604,	TEL. 312-663-9415
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Date Received for Review: Date Review Completed: 5/14
TO: Tim Boos/Phil Smith
FROM: Zena Gold-Kaufman FOX
SUBJECT: ChemetCO FILO523
Sample Description: Case # 2882
* Dioxin Samples
Project Data Status: COMPREC
NO hits for 2,3,7,8 TCOD Several hits in other isomers.

Additional Comments:

None

Book No. 6
Page No. 70

## UNITED STATES ENVIRONMENTAL PROTECTION AGENCY REGION V

ATE	5/12/87	
ECT.	Review of Region V CLP Data 5/1/27 Received for Review on	
ROM.	Review of Region V CLP Data 5/1/87  Received for Review on 5/1/87  Curtis Ross, Director (5SCRL) Latich & Chuida  Central Regional Laboratory  Data User: 50	
TO:	Data User:	<u>P</u> >
	We have reviewed the data for the following case(s).	
	SITE NAME: Chinictes SMD Case No. 345 2782E	
	SITE NAME: Chepictes— SMD Case No. 345 2782E  No. of D.U./Activity  EFA Cata Set No. 5F 3916 Samples: 5 Numbers 4651 072100	
•	WERL NO. 875805505 - 875805505	
	SMO Traffic No. E01-E05  CLP Laboratory: Intemple for Review: 7	
	Following are our findings.	
	1. CALIBRATION: ION RATIOS AND RESPONSE FACTORS ARE OK	
	2. COLUMN SEPARATION IS ACCEPTABLE	
	3. SURROGATE RECOVERY IS OK	
	4. MATRIX SPIKE AND DUPLICATE ARE IN AGREEMENT	
	- THE BLANK WAS CLEAN. DETECTION LIMITS ARE OK.	
, kl	6. SAMPLES ARE REPORTED ON A DRY WEIGHT BASIS. SPOT CHECK OF	N
	CALCULATIONS WAS OK. Patrick J. Chwille 5-12-87	
	5-/2-87	
	Data are acceptable for use.  Data are acceptable for use with qualifications noted above.  Data are preliminary - pending verification by Contractor Laboratory.  Data are unacceptable.	
	cc: Dr. Alfred Haeberer/Joan Fisk/Gary Ward, EPA Support Services Ross K. Robeson, EMSL-Las Vegas Don Trees, CLP/Sample Management Office	

## UNITED STATES ENVIRONMENTAL PROTECTION AGENCY REGION $\mathbf V$

## ESD/Central Regional Laboratory DATA TRACKING FORM FOR CONTRACT SAMPLES

## TRIANGLE LABORATORIES, INC.

919 544-5729

ED MAY 1 4 1927

CASE NARRATIVE

MAY 1 1987

DATE: April 25, 1987 CLIENT NO.: SAS 2882E TLI NO.: 8701364

Ole emiliario (g. 1903) 1883 Carrio (g. 1903) 18 (Charling Van) (g. 1903) OBJECTIVE: ANALYSIS OF SEDIMENT, SOIL AND SLAG SAMPLES FOR THE PRESENCE OF TETRA THROUGH OCTA CHLORO-DIBENZODIOXINS AND FURANS

The samples were extracted by the enclosed protocol. On the sample data sheets the concentration is given in parts per billion (ppb). "RT" is the retention time on the gas chromatographic column in minutes and seconds, "number" is number of isomers in the totals reported for each group, "ratio" is the ratio observed for the M to M+2 ions for tetra through penta chlorinated compounds and M+2 to M+4 for the hexa through octa. "DL" is the detection limit in parts per billion. For quantitation, the sum of the areas for the two masses monitored is used. When no peak is detected an area of 2 counts for each ion (total 4 counts) is used to calculate the detection limit.

The samples were spiked with 10 ng of 13C-2378-TCDD, 37C1-2378-TCDD, 13C-2378-TCDF, 13C-123478-HxCDF, 13C-12378-PCDD, 123678-HxCDD, 13C-1234678-HpCDD, and 20 ng of 13C-OCDD prior to the extraction. For GC/MS analysis, the final extract was dissolved in 20 ul of toluene containing 13C-1234-TCDD and 13C-123789-HxCDD at a concentration of 500 pg/ul to measure the recovery of the 13C- labeled internal standards.

Samples were analyzed using a VG 7070H mass spectrometer and 11-250 data system, operated in the selected ion recording mode, at a resolution of 5000. A Varian 3700 GC was employed, with a DB-5 60m x 0.32mm id fused silica capillary column. microliter of the 20 microliters final sample volume injected, splitless, at a column temperature of 150 deg C, heated ballistically to 190 deg C, then programmed at 3 deg/min to 300 deg C. A continuing calibration was demonstrated by injecting a solution of the analytes at a concentration of 100 pg/ul for the tetra isomers, 500 pg/ul for the penta though hepta and 1000 pg/ul for the octa, and a constant value of 100 pg/ul for 13C-2378-TCDD, 13C-2378-TCDF, 13C-123478-HxCDF, 37C1-2378-TCDD, 13C-12378-PCDD, 13C-123678-HxCDD, 13C-1234678-HpCDD, and 200 pg/ul 13C-OCDD. Response factors (RF) were calculated for the analytes from this continuing calibration. For the totals in each group the response factor is taken as the average of the response factors for the individual isomers listed. Note that in some cases, when only one isomer is present, there may be a discrepancy between the amount given for the individual isomer and the total since a different response factor may be used

relta RF on the und difference between the undiactor in the initial calibration.

Positive identification criteria for chlorinated and furans are as follows:

1) Ratio of M+ to M+2 or M+2 to M+4 is within 20% of the Control theoretical value, except for the tetrachloro which are taken within 13%.

The ranges of the ratios for identification of management of the compounds:

M+2/M+4 hexa 0.98-1.48hepta 0.82-1.24 octa 0.70-1.06

- 2) Retention time for analytes is within 3 seconds of the corresponding 13C internal standard or surrogate standard.
- 3) The identification of specific isomers that do not have corresponding 13C12-labeled standards is done by comparison of the retention time of the analyte to the nearest internal standard retention time with reference to the comparable retention times found in the continuing calibration.
- If the retention time and the ratio are correct for identification of an isomer, the signal to noise ratio (S:N) must be greater than 2.5.
- 5) For confirmation of 2378-TCDD and 2378-TCDF, the samples are run on a second GC column. 2378-TCDF is not fully resolved from other TCDF isomers on a DB-5 column, so the concentration obtained from the full screen analysis is considered a maximum. The concentration of 2378-TCDF obtained on the second column is sometimes less than that indicated by the full screen analysis, because of the increased GC resolution.
- For confirmation of 2378-TCDD and 2378-TCDF, resolution from its nearest isomer must be >25% valley. The ions for masses corresponding to M-COCl are also monitored for confirmations. The column performance check solution contains the following isomers: 2347, 2348, 2378, 2367, 3467, and 1368-TCDF, and 1478, 2378, 1237, 1238, and 1234-TCDD.

Sample E05 was used for the matrix spike and matrix spike duplicate. The matrix spike was 5 ng per sample of 2378-TCDD and 2378-TCDF. 25 ng for the penta though hepta chlorinated compounds, and 50 ng for the octa. Calculations for the percent recovery and precision are enclosed.

### GC/MS Conditions (FULL SCREEN ANALYSES)

Gas Chromatography:

Instrument: Varian 3700

Capillary Column: a. Manufacturer - J&W Scientific

b. Liquid phase - DB-5 Length - 60 m c. - 0.25 mm d. I.D

e. film thickness -0.25 microns

: Helium Carrier gas Head pressure : 28 psi

Flow thru column: 1 to 2 ml/min.

Injection type : Splitless for 30 sec.

Initial isothermal temperature :150 deg C for 30 sec.

Initial temperature program rate: to 190 deg C ballistically

Final temperature program rate : to 300 deg C @ 3 deg/min

### GC/MS Conditions (2378-TCDD CONFIRMATION)

Gas Chromatography:

Instument: Varian 3700

Column: a. Manufacturer - 1) Supelco, inc

2) J&W Scientific

- 1) SP-2331 Liquid phase

2) DB-17

-1) 30 m Length 2) 15 m

I.D - 0.32 mm d.

- 0.25 microns film thickness e.

Carrier gas : Helium Head pressure : 14 psi

Flow thru column: 1 to 2 ml/min.

Injection type : Splitless for 30 sec.

Initial isothermal temperature :150 deg C for 30 sec.

Initial temperature program rate: to 170 deg C ballistically Final temperature program rate : to 220 deg C @ 4 deg/min

### Mass Spectrometry:

متواداته.

Instrument: VG Micromass 7070H

Ionization mode : EI, positive ion

Reactant gas : N/A Resolution : 5000

Scan mode : selected ion recording

Switching mode : voltage Reference standard: PFK

### EXTRACTION OF SOIL AND ASH SAMPLES

After addition of an equal amount of anhydrous sodium sulfate and internal standard and surigate spike solution, the samples are extracted with foluene in a Soxhlet apparatus for 16 hours. The extract is chromatographed as described below.

### PURIFICATION

Column Al: Silica gel with basic and acidic layers. From the bottom, these layers are: I g activated silica gel, 2 g silica gel with 1N sodium hydroxide (2;1). I g silica gel, 12 g silica gel with concentrated sulfuric acid (2;2), 2 g silica gel and 1 cm of sodium sulfate on the top.

Column A2: Directly under column 1, a 2nd column of 6 g A-948 activated alumina with 10% water. The sample is loaded onto column 1 and the column is rinsed with 90 ml hexane. Column 1 is removed and the sample is eluted from column 2 with 20 ml of 1% methylene chloride in hexane (which is set aside for PCB analysis or in case of breakthrough) and 20 ml of 20% methylene chloride in hexane (dioxin fraction).

Column B: Mixed 124 g 545 celite and 10.7 g AX-21 carbon. Frewash with: 2 ml 50% benzene/ethyl acetate, 1 ml 50% methylene chloride/cyclohexane, and 2 ml hexane. After adding 1 ml of hexane, the dioxin fraction is loaded on the column, which is then rinsed with 2 ml 50% methylene chloride/hexane followed by 2 ml of 50% benzene/ethyl acetate. The column is inverted, and the sample is eluted with 4 ml of toluene.

The sample is evaporated to dryness with nitrogen and redissolved in 20 ul of toluene containing the recovery standards 13C12-1234-TCDD and 13C12-123789-HxCDD, for GC/MS analysis.

EXAMPLE CALCULATIONS

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SURROGATE, SAMPLE, OR INTERNAL STANDARD RESPONSE FACTORS:

(A<sub>S</sub>)(I<sub>iS</sub>)

= RF

$${A_{S}(I_{iS}) \atop --------} = RF {A_{IS}(I_{S})}$$

WHERE:

Single.

 $A_s$  = AREA OF SAMPLE <FROM RIC> 2297+2887 = 5184

A is = AREA OF APPROPRIATE INTERNAL STANDARD 2281+2764 = 5045

I is = CONCENTRATION OF INTERNAL STANDARD 100 Poffici

Is = CONCENTRATION OF ANALYTE 100 PS/kil

EXAMPLE: FROM FILE MAPOOTTO

The areas of the 2 masses monitored for the analyte are added and compared to the sum of the areas for the 2 masses monitored for the internal standard, except for the surrogate 37Cl-TCDD where only one mass is monitored (328), which is compared to the 334 ion in the internal standard, 13C12-2378-TCDD.

### SURROGATE \* RECOVERY:

1). Amount of surrogate found = 
$$\frac{(A_s)(I_{is})}{(A_{is})(RF)}$$

$$\beta C 12 - T c D F$$

WHERE:

TRE:
$$A_s = AREA OF SAMPLE 4336 + 5387 = 9723$$

A IS = AREA OF INTERNAL STANDARD
$$13C(2-2)75-700$$

$$1 = \text{AMOUNT OF INTERNAL STANDARD IN TOTAL EXTRACT} / 0.25$$

RF = RESPONSE FACTOR FOR SURROGATE (from Continuing Calibration 
$$Mf70770$$
) /.337

Recovery (for above sample) = 
$$\frac{[/0.44]}{[/0]} \times 100 = /04 \times \text{Recovery}$$

### CALCULATIONS OF INTERNAL STANDARD RECOVERY:

1).  $(A_{IS})(I_{RS})$ ----= Amount of Internal Standard Found

13012-2371-1200 WHERE:

 $A_{IS}$  = AREA FOUND FOR INTERNAL STANDARD 3/0/+3866 = 6967

 $A_{RS}$  = AREA FOUND FOR RECOVERY STANDARD 2572 + 3208 = 5720 (13C-1234-TCDD)

I<sub>RS</sub> = AMOUNT OF RECOVERY STANDARD IN EXTRACT FOR PS/FIL

RF<sub>IS</sub> = RESPONSE FACTOR FOR INTERNAL STANDARD 1.356

(from Continuing Calibration m \$7070)

13C12 2378-TCDD

FILE: B&7077/

(6967)(500 p)/we= 449.1 ps/fil

2). % Recovery = (Amount of Int. Std. Found) 100 (Amount of Int. Std. Added)

 $\frac{(449)}{(500)} \times 100 = 69.68 \text{ Recovery}$ 

CALCULATIONS OF DETECTION LIMITS:

DL: =  $(A_S)(I_{S}) \times 2.5$ (A<sub>IS</sub>)(WT.)(RF) RECEIVED MAY 1 4 1987

WHERE:

Ac = Area of sample channel integrated through the center of noise across a region corresponding to the baseline peak width of the standard compound (continuing calibration)

AIS = AREA OF INTERNAL STANDARD

13612 - 2378 - 7000 3101 + 3866 = 6967

IIS = AMOUNT OF INTERNAL STANDARD

2.5 = FACTOR FOR SIGNAL/NOISE ACCEPTANCE i.e.:  $s/n \ge 2.5$  for acceptance of signal statistically significant.

RF = RESPONSE FACTOR FOR ANALYTE 1.028(from Continuing Calibration  $M f7 \circ 720$ )

WT. = WEIGHT OF SAMPLE / 0.0 \$

EXAMPLE:

 $\frac{(32.8)(log)}{(6967)(log)(lo28)} \times 2.5 = .01 ppb$ 

\* 2 used for background (no peaks)

CALCULATIONS OF SAMPLE RESULT:

[A<sub>s</sub>][I<sub>is</sub>] concentration = [A<sub>is</sub>][RF<sub>s</sub>][W]

WHERE:

HERE:  $A_s$  = AREA FOUND FOR SAMPLE  $A_{is}$  = AREA FOUND FOR INTERNAL STANDARD 2277 + 26/9 = 4/9 L  $A_{is}$  = AMOUNT OF INT. STD. /0 ~8

RFs = RESPONSE FACTOR FOR ANALYTE 1.499
(from Continuing Calibration m870770)

W = MASS OF SAMPLE 7.03 3

EXAMPLE: 2378-TODF FILE: M870777

(10296)(10ng) (4896)(1.455)(7.035) = 1.996 ng/g = 1.996 pps

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TLI#8701364 DATE 4-24-87

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### TRIANGLE LABORATORIES, INC. MATRIX SPIKE

SAMPLE # SAS2882E05 MS

ISOMER 2378-TCDD	NATIVE (ppb)	SPIKED (ppb)	OBSERVED (ppb)	"RECOVERYCE TO THE TOTAL	
123478-PCDD	ND(.002)	2.43	2.48	102%	lo <sub>n</sub>
123478-HxCDD 123679-HxCDD 123789-HxCDD	ND(.002) ND(.002) ND(.002)	2.43 2.43 2.43	2.27 2.49 2.34	93% 102% 96%	<i>**</i> €>
1234678-HpCDD	ND(.003)	2.43	2.60	107%	
OCDD	. 06	4.87	4.91	100%	
2378-TCDF	ND(.001)	.49	. 55	113%	
12378-PCDF	ND(.001)	2.43	2.45	101%	
234678-PCDF	ND(.001)	2.43	2.43	100%	
123478-HXCDF 123678-HxCDF 234678-HXCDF 123789-HXCDF	ND(.001) ND(.001) ND(.001) ND(.001)	2.43 2.43 2.43 2.43	2.39 2.46 2.38 2.37	98% 101% 98% 97%	
1234678-HpCDF 1234789-HpCDF	.02 ND(.011)	2.43 2.43	2.53 2.73	103% 112%	
OCDF	. 05	4.87	5.18	105%	

TLI#8701364 DATE 4-24-87

## TRIANGLE LABORATORIES, INC. MATRIX SPIKE

SAMPLE # SAS2882E05 MSD

ISOMER	NATIVE (ppb)	SPIKED	OBSERVED	%RECOVERY
2378-TCDD	ND(.001)	. 49	.56	114%
123478-PCDD	ND(.002)	2.47	2.67	108%
123478-HxCDD 123679-HxCDD 123789-HxCDD	ND(.002) ND(.002) ND(.002)	2.47 2.47 2.47	2.15 2.44 2.36	87% 99% 96%
1234678-HpCDD	ND(.003)	2.47	2.70	109%
OCDD	. 06	4.94	5.12	102%
2378-TCDF	ND(.001)	. 49	.59	120%
12378-PCDF	ND(.001)	2.47	2.54	103%
234678-PCDF	ND(.001)	2.47	2.63	107%
123478-HXCDF 123678-HXCDF 234678-HXCDF 123789-HXCDF	ND(.001) ND(.001) ND(.001) ND(.001)	2.47 2.47 2.47 2.47	2.33 2.33 2.35 2.34	94% 94% 95% 95%
1234678-HpCDF 1234789-HpCDF	.02 ND(.011)	2.47 2.47	2.61 2.82	105% 114%
OCDF	. 05	4.94	5.28	106%

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# TRIANGLE LABORATORIES, INC. MATRIX SPIKE SAMPLE # SAS2882E05 MS AND SAS2882E05 MSD

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ISOMER	MS(TOTAL)	MSD(TOTAL)	AVERAGE (ng)	ABS	DEVIATION %DEV
2378-TCDD	5.402	5,657	<b>5.</b> 530	. 128	2%
TOTAL TODD	5.402	5.657		. 128	2%
123478-PCDD	25.470		26.220	.750	3%
TOTAL PCDD	25.470		26.220	. 750	3%
123478-HxCDD	23.282	21.768	22.525	757	-3%
123679-HxCDD	25.603	24.683	25.143	460	-2%
123789-HxCDD	24.011	23.883			-0%
TOTAL HXCDD	72.835	70.172	71.503	-1.331	-2%
1234678-HpCDD	26.661	27.324	26.992	. 332	1%
TOTAL HPCDD	26.661	27.324		. 332	1%
OCDD	50.374	51,814	51.094	.720	1%
				·	<del>-</del>
2378-TCDF	5.679	5.930	5.805	. 126	2%
TOTAL TODE	5.679	5.930	5.805	. 126	2%
12378-PCDF	25.172	25.745	25.459	. 287	1%
234678-PCDF	24.915	26.646	25.780	. 865	3%
TOTAL PCDF	50.138	52.209	51,174	1.035	2%
123478-HXCDF	24.556	23.539	24.047	508	-2%
123678-HxCDF	25.223	23.569	24.396	827	-3%
234678-HXCDF	24,443	23.802	24.122	-,320	-1%
123789-HXCDF	24.289	23.661	23.975	314	-1%
TOTAL HECDF	98.592	94.531	96.561	-2.031	-2%
1234678-HpCDF	25.963	26.383	26.173	.210	1%
1234789-HpCDF	28.068	28.518	28.293	.225	1%
TOTAL HPCDF	53.887	54.749	54.318	. 431	1%
OCDF	53.219	53,423	53.321	. 102	0%

## TRIANGLE LABORATORIES, INC. PCDD/PCDF ANALYSIS

ANALYST DATE SAMPLE WEIGHT SAMPLE ID	SAS 2882 TL		FILE # CONCAL # TLI #	870136	70
		NUMBER		RATIO	RT
2378-TCDD	ND		0.011	15.400	26.36
TOTAL TODD	ND	0	0.011	15.400	
12378-PCDD	ND		0.015		32.30
TOTAL PODD	ND	0	0.015		
123478-HxCDD	ND		0.002	1.000	
123678-HxCDD	ND		0.002	1.000	
123789-HxCDD	ND		0.003	1.000	
TOTAL HXCDD	ND	0	0.025	1.241	
1234678-HpCDD	ND		0.003	1.000	
TOTAL HPCDD	ND	0	0.003	1.000	
OCDD	ND		0.021	0.100	49.58
2378-TCDF	ND		0.023	0.626	25.54
TOTAL TODE	ND	0	0.023	0.626	
12379-PCDF	ND		0.001	1.000	
23478-PCDF	ND		0.040	0.850	32.08
TOTAL PCDF	ND	0	0.032	0.850	
123478-HxCDF	ND		0.001	1.000	
123678-HxCDF	ND		0.001	1.000	
234678-HxCDF	ND		0.001	1.000	
123789-HxCDF	ND		0.058	1.613	38.00
TOTAL HxCDF	ND	0	0.052	1.613	
1234678-HpCDF	ND		0.002	1.000	
1234789-HpCDF	ND		0.002	1.000	
TOTAL HpCDF	ND	0	0.002	1.000	
OCDF	ND		0.003	1.000	
SURROGATE RESU	LTS SUMMARY				
NAME	CONC. (ppb)	% RECOVERY	RATIO	RT	==
13C12-TCDF	1.04	104.38	0.805	25.56	
37Cl-TCDD	1.13			26.37	
13C12-HxCDF			1.218		
INTERNAL STAND					<del>-</del> -
NAME		% RECOVERY		RT	
2378-13C12-TCD	D 0.90	89.82	0.802	26.36	- <del>-</del>
13C12-PCDD	0.88	88.33	0.643		
13C12-HxCDD		95.42	1.296		
13C12-HpCDD		83.16	1.105		
13C12-OCDD	1.53	76.60	0.905		

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## TRIANGLE LABORATORIES, INC. PODD/PODF ANALYSIS

ANALYST DATE SAMPLE WEIGHT SAMPLE ID			FILE # CONCAL # TLI #		(-)
NAME	CONC. (ppb)	NUMBER	DL DL	RATIO	ET
2378-TCDD	ND		0.157	0.577	26.34
TOTAL TOPD	15.108	ප		0.794	
12378-PODD				0.539	32,32
TOTAL PODD	12.905	11		0.603	
123478-HxCDD	0.511			1.301	<b>3</b> 7.03
123673-HxCDD	1.173			1.420	37.10
123789-HxCDD	1.812			1.177	37.37
TOTAL HxCDD		7		1.267	
1234678-HpCDD	10.743			1.050	43.50
_ <u>_</u>	20.955	2		1.027	
CCDD	20.527			0.874	49.59
2378-TCDF	15.972			0.750	25.53
TOTAL TCDF	71.147	17		0.750	
12378-PCDF	2.103			0.659	31,13
23478-PCDF	6.193			0.617	32;06
TOTAL PCDF	48.197	14		0.629	
123478-HxCDF	10.676			1.231	35.57
123678-HxCDF	3.744			1.217	<b>3</b> 6.06
234678-HxCDF	8.475			1.236	36.52
123789-HxCDF	0.675			1.188	37.58
TOTAL HXCDF		12		1.228	
1234678-HpCDF				1.007	42.22
1234789-HpCDF				0.972	44.28
TOTAL HPCDF	58.975	4		1.002	
OCDF	64.718			0.896	50.12

### SURROGATE RESULTS SUMMARY

16(80)P

===========	=========			=========
NAME	CONC. (ppb)	% RECOVERY	RATIO	RT
13C12-TCDF 37C1-TCDD 13C12-HxCDF *	1.953 1.998 2.238	107.05 109.51 122.63	0.788 0.761	25.52 26.34 35.58

### INTERNAL STANDARDS RECOVERY RESULTS

==============	=========	===========	:========	========
NAME	CONC. (ppb)	% RECOVERY	RATIO	RT
2378-13C12-TCDD 13C12-PCDD 13C12-HxCDD 13C12-HpCDD 13C12-OCDD	1.613 1.474 1.670 1.467 2.626	88.42 80.78 91.49 80.37 71.95	0.847 0.722 1.223 1.149 0.848	26.33 32.31 37.10 43.50 49.59

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## TRIANGLE LABORATORIES, INC 2,3,7,8-TCDD/TCDF ANALYSIS

SAMPLE WEIGHT	MDC 4/25/87 5.48 SAS 2882E01	FILE # CONCAL # TLI #	M87079 M87079 870136	)5
NAME	CONC (ng/g)	DL	RATIO	RT
2378-TCDF 2278-TCDD	2.55 0.11		0.814 0.838	
SURROGATE RESULT	rs summary			
NAME	CONC (ng/g)	% RECOVERY	RATIO	RT
'3C12-TCDF	2 11	115.42	0.804	27.24
m37C1-TCDD	2.27			22.55
INTERNAL STANDA	2.27	124.54		
INTERNAL STANDA	2.27	124.54 ESULTS		

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## TRIANGLE LABORATORIES, INC. PCDD/PCDF ANALYSIS

DATE SAMPLE WEIGHT			FILE # CONCAL # TLI #		<b>'</b> O
	SAS 2882 E02				
	CONC. (ppb)				
2378-TCDD	ND	2	0.034		
TOTAL TODD	0.632	2		0.750	
12378-PCDD				0.542	32.30
TOTAL PODD	0.989	7		0.600	
123478-HxCDD	0.063			1.051	37.03
123678-HxCDD	0.152			1.000	37,11
123789-HmCDD	0.226			1.182	37.40
TOTAL HxCDD		6		1.220	
1234678-HpCDD TOTAL HpCDD OCDD	1.678			1.084	43.51
TOTAL HPODD	3.269	2		1.056	
OCDD	5.053			0.873	50.00
2378-TCDF	1.996			0.737	25, 53
TOTAL TCDF		13		0.738	
12378-PCDF	0.259			0.600	31.13
23478-PCDF				0.597	32.05
TOTAL PCDF		11		0.608	
123478-HxCDF	1.426			1.396	
123678-HxCDF				1.429	
234678-HxCDF	1.171			1.223	
123789-HxCDF			·	1.296	38.00
TOTAL HMCDF		10		1.305	
1234678-HpCDF	4.378			1.018	42.23
1234789-HpCDF	1.077			0.995	44.29
TOTAL HPCDF	8.406	4		1.010	
OCDF	9.532			0.888	50.12
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### SURROGATE RESULTS SUMMARY

14(B)14

NAME	CONC. (ppb)	% RECOVERY	RATIO	RT
13C12-TCDF 37C1-TCDD 13C12-HxCDF	1.399 1.379 1.426	98.32 96.97 100.24	0.807	25.51 26.33 35.58
TOCIZ TRODI	1.420	100.24	1.219	00.00

### INTERNAL STANDARDS RECOVERY RESULTS

· NAME	CONC. (ppb)	%	RECOVERY	RATIO	RT
2378-13C12-TCDI	1.307		91.90	0.869	26.32
13C12-PCDD	1.237		31.90 36.99	0.647	32.30
13C12-HxCDD	1.324		93.07	1.241	37.10
13C12-HpCDD	1.204		84.62	1.063	43.51
13C12-OCDD	2.250		79.10	0.878	49.59

## TRIANGLE LABORATORIES, INC 2,3,7,8-TCDD/TCDF ANALYSIS

ANALYST DATE SAMPLE WEIGHT SAMPLE ID		FILE # CONCAL # TLI #		15
, NAME	CONC (ng/g)	DL	RATIO	RT
2378-TODF 2375-TODD			0.802 0.683	
SURFOGATE RESULI	S SUMMARY			
	============	==========	:=======	=======
NAME	CONC (rg/g)		RATIO	RT
13C12-TCDF	CONC (ng/g)	% RECOVERY 101.72	0.799	
13C12-TCDF	CONC (ng/g) 1.45 1.55	% RECOVERY 101.72 109.19	0.799	26.47
13C12-TCDF 37C1-TCDD INTERNAL STANDAR	CONC (ng/g) 1.45 1.55	% RECOVERY  101.72 109.19  ESULTS	0.799	26.47

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## TRIANGLE LABORATORIES, INC. PCDD/PCDF ANALYSIS

ANALYST	JAJ		FILE #	M37077	15
	4-23-87		CONCAL #	M87077	Ģ
SAMPLE WEIGHT			TLI #	870136	-1
SAMPLE ID	SAS 2882 E03				
NAME		=========	=======================================	=======	======
NAME	CONC. (ppb)	NUMBER	DL	RATIO	EL
2378-TCDD	ND	б	0.021	0.822	26 36
TOTAL TOPD	0.297	6		0.794	
12378-PCDD				0.543	32,35
TOTAL PODD	0.435	7		0.586	
123478-HxCDD	0.029			1.195	37.07
123678-HMCDD	0.091			1.132	
123789-HmCDD	0.140			1.322	37.43
TOTAL HMCDD	0.988	6		1.338	
1234678-HpCDD	1.080			1.024	43.55
TOTAL HPCDD	2.071	2		1.015	
OCDD	4.291			0.882	50,04
2378-TCDF	0.659			0.772	25.53
TOTAL TCDF	3.151	15		0.749	
12378-PCDF	0.091			0.551	31.10
23478-PCDF	0.250			0.635	<b>3</b> 2.08
TOTAL PCDF	2.184	13		0.601	i
123478-HxCDF				1.274	36.01
123678-HxCDF				1.442	36.11
234678-HxCDF				1.280	36.56
123789-HxCDF			0.001	1.000	
TOTAL HXCDF	2.560	10		1.268	
1234678-HpCDF	1.991			0.988	<b>4</b> 2.28
1234678-HpCDF 1234739-HpCDF	0.349			1.042	44.33
TOTAL HPCDF OCDF	3.411	4		0.989	
OCDF	3.281			0.892	<b>5</b> 0.16

### SURROGATE RESULTS SUMMARY

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NAME	CONC. (ppb)	% RECOVERY	RATIO	RT
13C12-TCDF 37C1-TCDD 13C12-HxCDF	1.198 1.263 1.095	106.98 112.75 97.75	0.762 1.193	25.53 26.36 36.02

### INTERNAL STANDARDS RECOVERY RESULTS

NAME C	CONC. (ppb)	% RECOVERY	RATIO	RT
2378-13C12-TCDD	1.027	91.71	0.804	26.35
13C12-PCDD	0.947	84.53	0.606 1.050	32.34 37.14
13C12-HxCDD	1.040 1.090	92.89 97.31	1.250 1.042	37.14 43.54
13C12-HpCDD 13C12-OCDD	2.200	98.23	0.898	50.03